AMENDMENTS TO THE CLAIMS

Claims 1-6 (cancelled)

- 7. (currently amended) A method of detecting a first target sequence comprising a first target domain, a second adjacent target domain and a <u>single stranded</u> poly(A) sequence, said method comprising:
 - a) hybridizing a first probe comprising:
 - i) an upstream universal priming site (UUP); and
 - ii) a first target-specific sequence substantially complementary to said first target domain; to said first target domain:
 - b) hybridizing a second probe comprising:
 - iii) a second target-specific sequence substantially complementary to said second <u>adjacent</u> target domain;
 - iv) a downstream universal priming site (DUP); to said second adjacent target domain;

wherein at least one of said first and second probes comprises at least a first adapter sequence; , said poly (A) sequence remains single-stranded, and wherein said target sequence and said first and second probes form a ligation complex;

- c) contacting said ligation complex with a ligase to form a ligated complex;
- d) contacting said ligated complex with a support comprising a poly(T) sequence, such that said single stranded poly(A) sequence hybridizes with said poly(T) sequence;
- e) removing unhybridized first and second probe sequences;
- f) denaturing said ligation complex;
- g) amplifying the ligated first and second probes to generate a plurality of amplicons;
- h) contacting said amplicons with an array of capture probes to form assay complexes; and
- i) detecting said assay complexes.
- 8. (original) A method according to claim 7 wherein said first target domain and said second target domain are directly adjacent.

- 9. (original) A method according to claim 7 wherein said first target domain and said second target domain are separated by at least one base and said method further includes contacting said ligation complex with a polymerase and at least one dNTP.
- 10. (previously amended) A method according to claim 7, 8 or 9 wherein one of said first and second probes comprises a label.
- 11. (original) A method according to claim 10 wherein said label is a primary label.
- 12. (original) A method according to claim 11 wherein said label is a fluorescent label.
- 13-14. (withdrawn)
- 15. (previously amended) A method according to claim 7, 8 or 9 wherein said amplifying is done by:
 - a) hybridizing a first universal primer to said UUP;
 - b) providing a polymerase and dNTPs such that said first universal primer is extended;
 - c) hybridizing a second universal primer to said DUP;
 - d) providing a polymerase and dNTPs such that said second universal primer is extended; and
 - e) repeating steps a) through d).
- 16. (previously amended) A method according to claim 7 wherein said array comprises:
 - a) a substrate with a patterned surface comprising discrete sites; and
 - b) a population of microspheres comprising at least a first subpopulation comprising a first capture probe and a second subpopulation comprising a second capture probe.
- 17.(original) A method according to claim 16 wherein said discrete sites comprise wells.
- 18. (original) A method according to claim 16 wherein said substrate comprises a fiber optic bundle.
- 19. (currently amended) A method according to claim 7, 8 or 9 wherein said support comprising a poly(T) sequence comprises magnetic beads comprising a poly(T) sequence.
- 20. (previously added) A method according to claim 15 wherein at least one of said first universal primers and said second universal primer comprises a label.